

Bats and human blood project: median cell sizes

prepare

```
pathWD <- "~/Documents/08_Pojekte/01_dRT-DC/220301_Bob_DataAnalysis_BatsHumans/20230829_AnimalTypeTempS  
setwd(pathWD)
```

load data space image

```
load(file.path(pathWD, paste0("RDataSpace_v8", ".RData")))
```

find median cell sizes and corresponding cells

```
for (iter.AnimalType in levels.AnimalType) {  
  print(paste0("Species = ", iter.AnimalType, ":"))  
  tbl.AnimalType <- combTable[combTable$animalType == iter.AnimalType, ]  
  medianArea <- median(tbl.AnimalType$area)  
  print(paste0("Median cell size = ", medianArea, " um^2"))  
  print("Cells:")  
  tbl.medAreaCells <- tbl.AnimalType[tbl.AnimalType$area == medianArea, c("dateCode", "tempLevel", "measN  
  print(tbl.medAreaCells)  
  write.csv(tbl.medAreaCells, file.path(pathWD, paste0("tbl", "_medianArea", iter.AnimalType, ".csv")),  
  print("")  
}
```

```
## [1] "Species = Human:"  
## [1] "Median cell size = 36.5585 um^2"  
## [1] "Cells:"  
## # A tibble: 141 x 12  
##   dateCode tempLevel measNum cellNum area velocity tau1 tau2 deltaDefoTau1  
##   <dbl> <fct> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>  
## 1 191108 Cold 2 109 36.6 4.15 4.94 0.883 0.135  
## 2 191108 Cold 2 948 36.6 4.01 4.43 1.87 0.159  
## 3 191108 Cold 3 1275 36.6 3.93 10.5 1.99 0.171  
## 4 191108 Cold 4 572 36.6 3.85 9.77 4.08 0.217  
## 5 191108 Cold 4 985 36.6 3.82 5.69 1.81 0.185  
## 6 191108 Cold 5 250 36.6 3.84 10.8 2.66 0.252  
## 7 191108 Cold 6 181 36.6 3.86 4.09 0.877 0.172  
## 8 191108 Cold 6 984 36.6 3.89 9.41 1.13 0.241  
## 9 191108 RT 2 255 36.6 4.23 7.31 1.12 0.222  
## 10 191108 RT 2 1117 36.6 4.17 7.82 0.864 0.238  
## # i 131 more rows  
## # i 3 more variables: deltaDefoTau2 <dbl>, eModulus <dbl>, viscosity <dbl>  
## [1] ""  
## [1] "Species = RousettusAegyptiacus:"
```

```

## [1] "Median cell size = 21.2704 um^2"
## [1] "Cells:"
## # A tibble: 1,281 x 12
##   dateCode tempLevel measNum cellNum area velocity tau1 tau2 deltaDefoTau1
##   <dbl> <fct>      <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
## 1 210812 Cold         1    314 21.3  3.74  4.15  2.37  0.118
## 2 210812 Cold         1    427 21.3  3.75  4.36  5.03  0.134
## 3 210812 Cold         1    503 21.3  3.75  4.67  1.71  0.137
## 4 210812 Cold         1    554 21.3  3.74  7.21  1.54  0.182
## 5 210812 Cold         1    778 21.3  3.75  3.33  1.69  0.126
## 6 210812 Cold         1    843 21.3  3.76  3.78  3.15  0.135
## 7 210812 Cold         1   2811 21.3  3.72 NaN    2.35    NaN
## 8 210812 Cold         1   3047 21.3  3.71  6.97  3.12  0.196
## 9 210812 Cold         1   3274 21.3  3.72  5.21  3.51  0.157
## 10 210812 Cold         1   3638 21.3  3.74  4.42  3.33  0.155
## # i 1,271 more rows
## # i 3 more variables: deltaDefoTau2 <dbl>, eModulus <dbl>, viscosity <dbl>
## [1] ""
## [1] "Species = NyctalusNoctula:"
## [1] "Median cell size = 19.3341 um^2"
## [1] "Cells:"
## # A tibble: 1,061 x 12
##   dateCode tempLevel measNum cellNum area velocity tau1 tau2 deltaDefoTau1
##   <dbl> <fct>      <dbl> <dbl> <dbl> <dbl> <dbl> <dbl> <dbl>
## 1 191015 Cold         2    778 19.3  4.07 NaN    4.17    NaN
## 2 191015 Cold         4     89 19.3  3.85  2.03 NaN    0.0619
## 3 191015 Cold         5    127 19.3  4.00  2.27 NaN    0.0729
## 4 191015 Cold         6    646 19.3  3.88  8.26 NaN    0.164
## 5 191015 Cold         7    155 19.3  3.93 NaN    4.00    NaN
## 6 191015 Cold        10     95 19.3  3.94 10.5  1.85  0.243
## 7 191015 Cold        10    263 19.3  3.96  6.58  2.61  0.192
## 8 191015 Cold        10    267 19.3  3.96 NaN    1.92    NaN
## 9 191015 RT           3    228 19.3  4.30 NaN    5.99    NaN
## 10 191015 RT           3    517 19.3  4.27  2.95 NaN    0.142
## # i 1,051 more rows
## # i 3 more variables: deltaDefoTau2 <dbl>, eModulus <dbl>, viscosity <dbl>
## [1] ""

```

```
rm(tbl.AnimalType,medianArea)
```